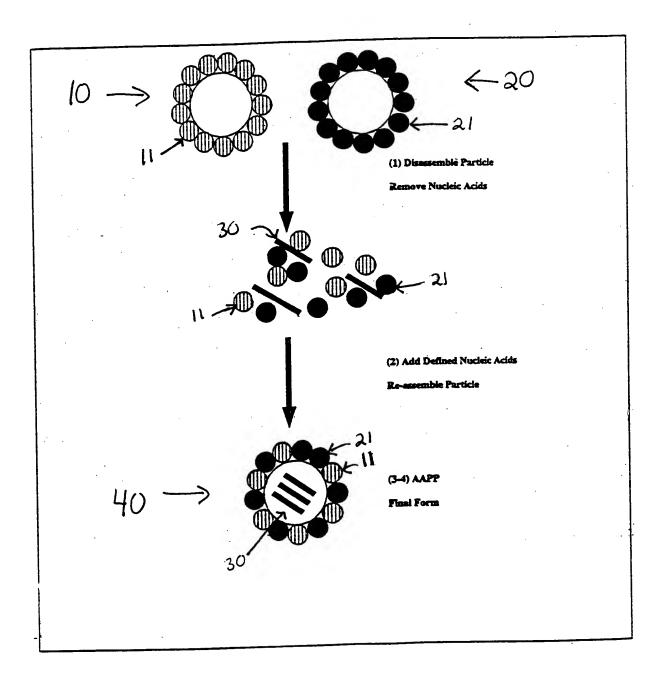
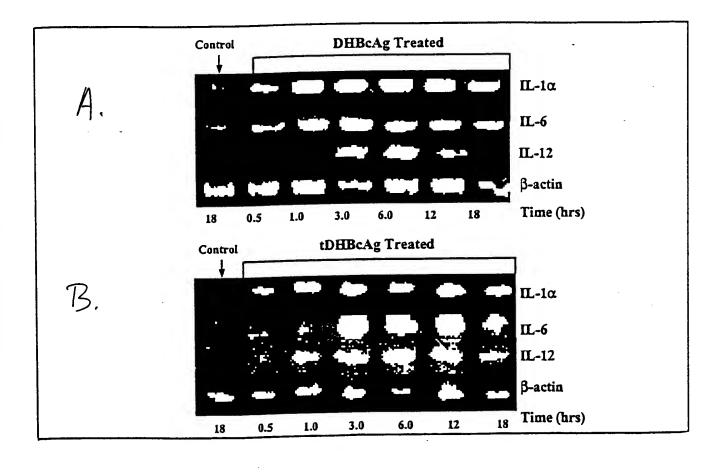
Amino Acid Sequence of Duck Hepatitis B Core Protein

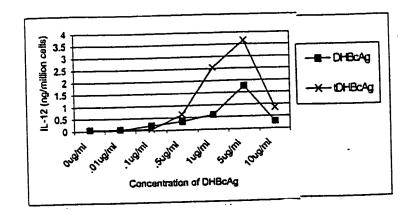
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Nucleic Acid Sequence of Duck Hepatitis B Core Protein

150 CAATAAAGAAACATGTTTTAATTGCAACTCACTTTGTGGATCTTATTGAAGACTTCTGGCAA 200 $\begin{smallmatrix} 100\\ \text{AAAAATTGATGATCTTGTAAGGGATGCGAAGGATGCTTTAGAACCTTATTGGAGATCAGATT} \end{smallmatrix}$ ATGGATATCAATGCTTCTAGAGCCTTAGCCAATGTTTATGATTTGCCAGATGATTTCTTCCC 250 ACCAGTTCCCGCAGGATATCTGATTCAGCACGAAGAGGCTGAGGAGATTCCTCTGGGAGATT ACTACTCAGGGTATGCATGAAATAGCTGAAGCCTTAAGAGCAGTTATACCACCTACTACAAC 350 TATTTAAACATCAGGAAGAAAGGATAGTTAGTTTCCAACCGGATTATCCTATTACTGCACGA $600\\ GAAGCAATCACTAGACCAATCCAAGTGGCTCAGGGAGGCAGAAAAACATCTTCGGGTACTAG$ CTCĞĞČTTCGCACTTGGCTATCAACACCTGAGAGATACAGAGGCCGAGATGCCČČÄACCATT 450 ATTGCTTTGGTGGCATTACAATTGTTTACTGTGGGGAGAAGCTAACGTTACTAATTATATTT AAAACCTCGTGGACTCGAACCTAGAAGAAGAAGTTAAAACCACAGTTGTCTATGGGAGAA 750 CGTAGTTCGAGCAGCACAGAAGATCTCCCTCGCCTAGGAAA

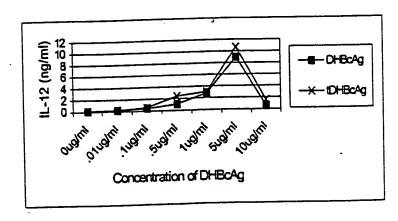


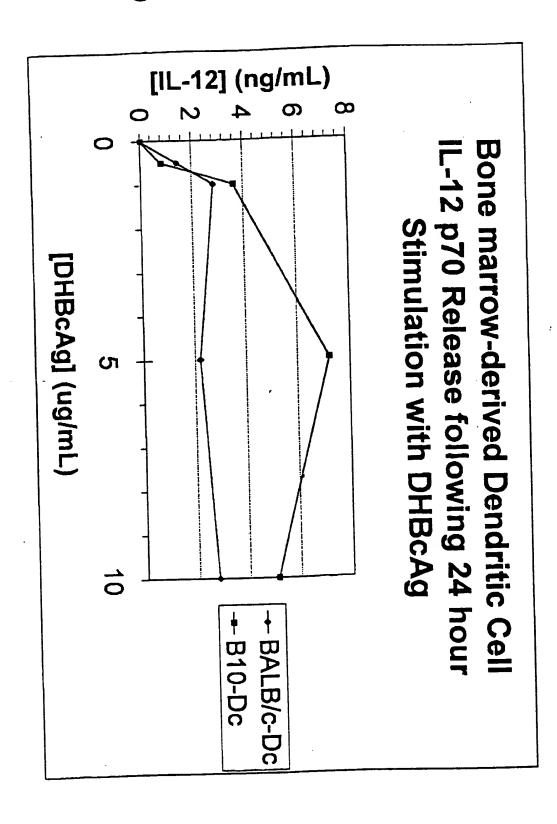


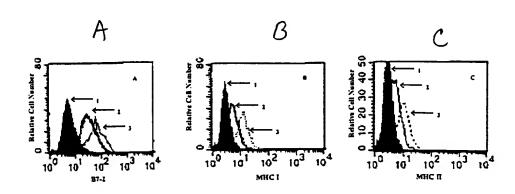


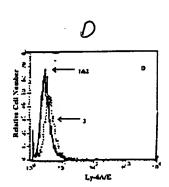
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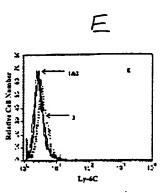
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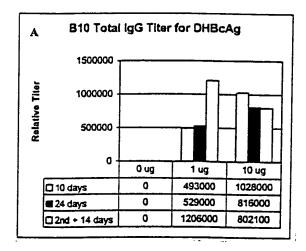




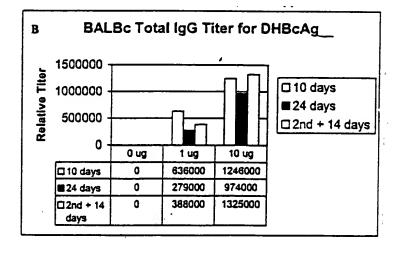


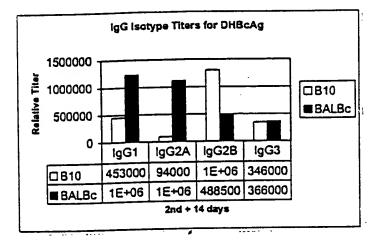


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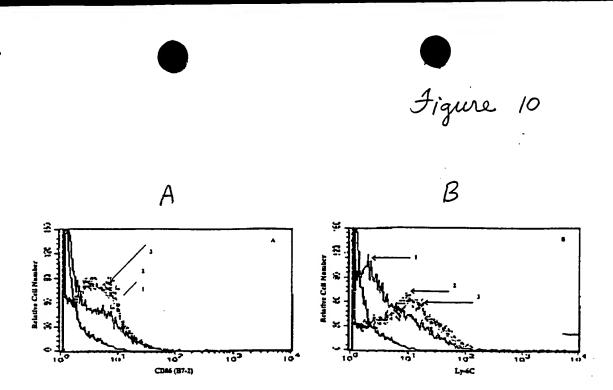


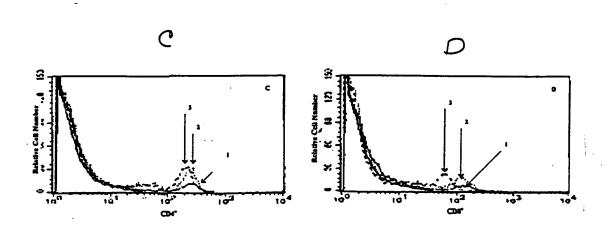
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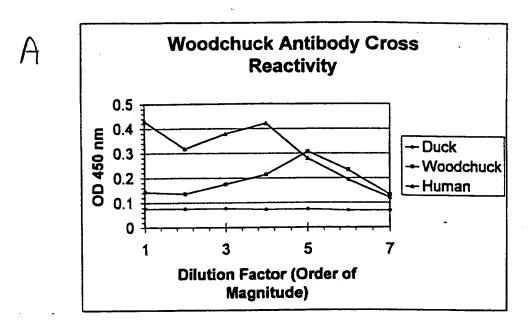


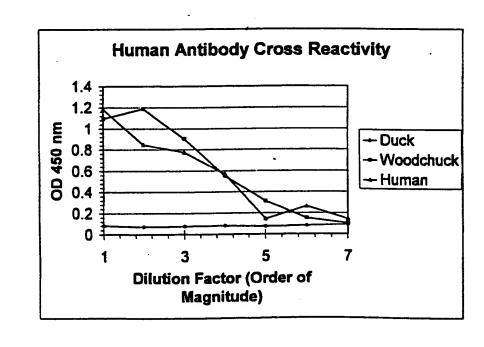


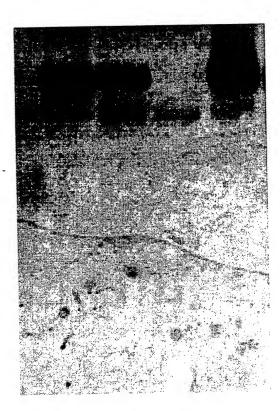
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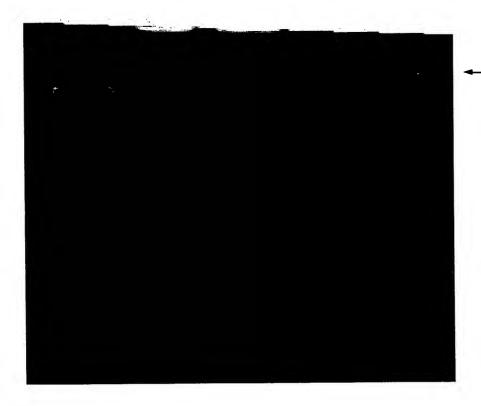






Duck HBcAg

1 2 3 4 5

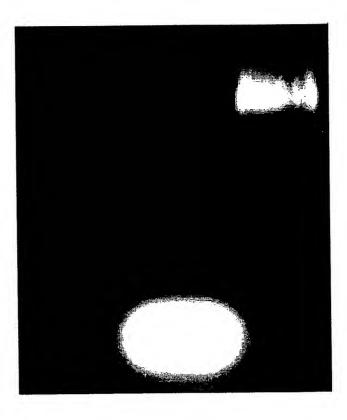


Duck HBcAg

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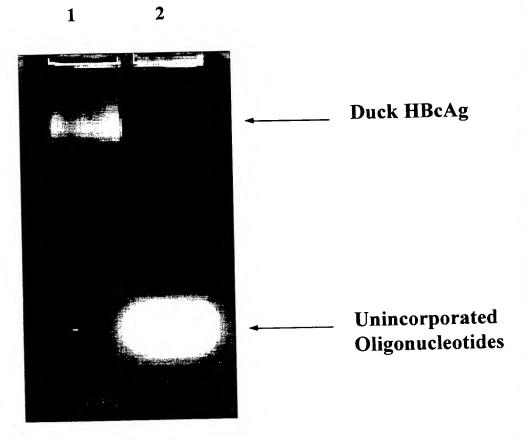
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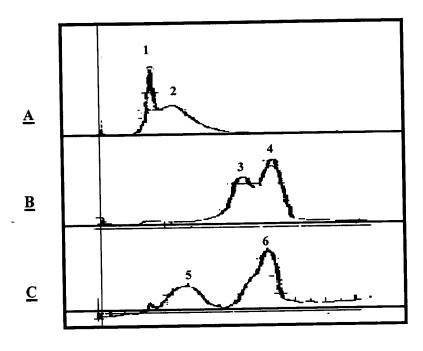


Duck HBcAg

Unincorporated Oligonucleotides







Amino Acid Sequence of Duck Hepatitis B Core Protein Mutant 1-239

239 RRAPS $^{1}_{\texttt{MDINASRALANVYDLPDDFFPKIDDLVRDAKDALEPYWR}}$ SDSIKKHVLIATHFVDLIEDFWQTTQGMHEIAEALRAVI 200 AQGGRKTSSGTRKPRGLEPRRRKVKTTVVYGRRRSKSRD PPTTTPVPAGYLIQHEEAEEIPLGDLFKHQEERIVSFQP GEANVTNYISRLRTWLSTPERYRGRDAPTIEAITRPIQV DYPITARIHAHLKAYAKINEESLDRARRLLWWHYNCLLW

Nucleic Acid Sequence of Duck Hepatitis B Core Protein Mutant 1-239

150 CAATAAAGAAACATGTTTAATTGCAACTCACTTTGTGGATCTTATTGAAGACTTCTGGCAA $\begin{smallmatrix} 100\\ \text{AAAAATTGATGATCTTGTAAGGGATGCGAAGGATGCTTTAGAACCTTATTGGAGATCAGATT} \end{smallmatrix}$ 50 ATGGATATCAATGCTTCTAGAGCCTTAGCCAATGTTTATGATTTGCCAGATGATTTCTTCCC CTCGGCTTCGCACTTGGCTATCAACACCTGAGAGATACAGAGGCCGAGATGCCCCAACCATT 200 ACTACTCAGGGTATGCATGAAATAGCTGAAGCCTTAAGAGCAGTTATACCACCTACTACAAC 600 GAAGCAATCACTAGACCAATCCAAGTGGCTCAGGGAGGCAGAAAAACATCTTCGGGTACTAG 450 ATTGCTTTGGTGGCATTACAATTGTTTACTGTGGGGAGAAGCTAACGTTACTAATTATATTT 718 GACGTTCAAAGTCCAGGGATAGGAGAGCCCCTTCA $\verb|AAAACCTCGTGGACTCGAACCTAGAAGAAGAAGAAGTTAAAACCACAGTTGTCTATGGGAGAA|$ 350 TATTTAAACATCAGGAAGAAAGGATAGTTAGTTTCCAACCGGATTATCCTATTACTGCACGA ACCAGTTCCCGCAGGATATCTGATTCAGCACGAAGAGGCTGAGGAGATTCCTCTGGGAGATT

Amino Acid Sequence of Truncated Duck Hepatitis B Core Protein

AQGGRKTSSGTRKPRGLEP GEANVTNYISRLRTWLSTPERYRGRDAPTIEAITRPIQV SDSIKKHVLIATHFVDLIEDFWQTTQGMHEIAEALRAVI MDINASRALANVYDLPDDFFPKIDDLVRDAKDALEPYWR DYPITARIHAHLKAYAKINEESLDRARRLLWWHYNCLLW PPTTTPVPAGYLIQHEEAEEIPLGDLFKHQEERIVSFQP

Nucleic Acid Sequence of Truncated Duck Hepatitis B Core Protein

150 CAATAAAGAAACATGTTTAATTGCAACTCACTTTGTGGATCTTATTGAAGACTTCTGGCAA 200 50 ATGGATATCAATGCTTCTAGAGCCTTAGCCAATGTTTATGATTTGCCAGATGATTTCTTCCC 400 ATTCATGCACACCTGAAAGCTTATGCAAAGATTAACGAGGAATCACTGGATAGGGCTAGGAG ${\tt AAAAATTGATGATCTTGTAAGGGATGCGAAGGATGCTTTAGAACCTTATTGGAGATCAGATT}$ $600\\ GAAGCAATCACTAGACCAATCCAAGTGGCTCAGGGAGGCAGAAAAACATCTTCGGGTACTAG$ CTCĞĞCTTCGCACTTGGCTATCAACACCTGAGAGATACAGAGGCCGAGATGCCCĞĞACCATT ACCAGTTCCCGCAGGATATCTGATTCAGCACGAAGAGGCTGAGGAGATTCCTCTGGGAGATT ACTACTCAGGGTATGCATGAAATAGCTGAAGCCTTAAGAGCAGTTATACCACCTACTACAAC 642 AAAACCTCGTGGACTCGAACCT ATTGCTTTGGTGGCATTACAATTGTTTACTGTGGGGAGAAGCTAACGTTACTAATTATATTT TATTTAAACATCAGGAAGAAAGGATAGTTAGTTTCCAACCGGATTATCCTATTACTGCACGA